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 Attorney's Docket No.: 11926

 Serial No. : 09/963,333
 015002 / 0017.CIP3DIV1

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In the Specification:

Replace the Sequence Listing with the enclosed, revised Sequence Listing.

Replace Table 10 beginning at page 171, with the following table:

Variance Table

Hugo	GID	MIMO	ID	VGX	Symbol	Desc	ription	
	iance Start		iance					
U73338	U73338	15657	0	GEN-	-69	Meth	nionine	
Synthase (SEQ								
	194		01)C>G		5 '			
	284	(-1)	11)C>T	1	5 '	ı		
	136		742G>P		V248N	1		
	252		858C>I	1	Silent	-		
	334		940G>A		D314N	1		
	699	1	305T>C	<u>.</u>	Silent	:		
	150	2	756A>G	;	D9190	3		
	207		813G>T		S938]	Ī.		
	209	2	815G>C	,	G939F	₹		
	444	5	050C>A	L	3 '			
	551	5	157G>A		3 '			
	573	5	179C>T	1	3 '			
	659	5	265T>C		3'			
	678	5	284T>C		3 '			
	874	5	480C>T		3 '			
	934		540A>G		3 '			
D78586	D78586	11401	0	GEN-	BR	CAD	PROTEIN	(SEQ
ID NO:2)								
	434		408C>T		Silent			
	313		287T>C		Silent			
	799		773A>G		Silent	•		
	255		229C>T		Silent			
	455	5	429G>A		R1810Ç)		
	507	5	481T>C		Silent			
	810		784C>T		Silent	•		
	128	6	102C>T		Silent	· ·		
	626		600C>T		Silent			
	686		660C>T		Silent			
U09178 U09178 274270 GEN-HA								
Dihydropyrimic		enase						
•	166		85T>C		C29R			

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577	496A>G	M166V	
638	557A>G		
1708	1627A>G		
3432	3351T>C	= :	
3682	3601C>T		
3730	3649G>A	_	
3925	3844A>G	3'	
3937	3856T>C	3'	
U19720 U19720		_	olate
Transporter (SLC19A1)			Jidee
175	80G>A	R27H	
341	246C>G		
791	696C>T		
1067	972G>A		
1337	1242C>A		
1997	1902T>C	3'	
2100	2005^2006insG	3'	
2582	2487T>G	3 '	
2617	2522C>T	3'	
2652	2557T>C	3'	
	0424 GEN-LUK	-	
folate carrier (RFC1)			ens reduced
431	431A>G		DEQ ID NO:5)
441	441A>G		
498	498C>T		
579	579G>C		
599	599G>C		
X02308 X02308		Intron EN-KL Th	
synthetase (SEQ ID NO:		EN-KT IL	ymidylate
1066	961T>C		
1136	1031A>G	3'	
1497	1392T>A	3' 3'	
		_	11
synthase, promoter (SE			iyıate
276			
321	276C>T	Intron	
452	321T>C	Intron	
457	452G>A	Intron	
491	457^insC	Intron	
533	491C>A	Intron	
624	533T>C	Intron	
	624A>C	Intron	
639	639A>G	Intron	
655	655T>C	Intron	

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D0059	96 D00596	5 188350 GEN-L	תוז.	Ното с	aniona
		synthase, exons 1, 2	3 1	Homo s	aprens
complete	cds (SEO ID	NO:8) (SEQ ID NO:3)	, ,, 4,	5, 6,	<i>'</i> ,
•	701	701A>C	Intron		
	716	716A>G	Intron		
	732	732T>C	Intron		
	1293	1293A>G	Intron		
	1322	1322C>G	Intron		
	1379	1379T>C	Intron		
	1590	1590C>T	Intron		
	1688	1688C>G	Intron		
	2401	2401A>G			
	2429	2429G>A	Intron Intron		
	2488	2423G>T	Intron		
	2594	2594G>T			
	2618	2618G>A	Intron		
	3083	3083G>A	Intron Intron		
	3125	3125G>A			
	3212	3212C>T	Intron		
	3619	3619T>A	Intron		
	3635	3635G>A	Intron		
	4256	4256G>A	Intron		
	4898	4898A>G	Intron		
	5006	5006C>T	Intron		
	5062	5062G>A	Intron		
	5167	5167G>A	Intron		
	11069		Intron		
	11238	11069A>G 11238C>T	Intron		
	11293		Intron		
	11422	11293T>G	Intron		
	11686	11422T>C	Intron		
	12598	11686C>T 12598T>C	Intron		
	13171	13171T>C	Intron		
	13298		Intron		
	13645	13298G>A	Intron		
	13751	13645T>C	Intron		
	13782	13751C>A	Intron		
	13806	13782T>C 13806T>C	Intron		
	13813		Intron		
	14479	13813T>C	Intron		
	14546	14479A>G	Intron		
	14585	14546^insT	Intron		
		14585C>T	Intron		
	14729 14787	14729G>A	Intron		
	14707	14787C>T	Intron		

14795G>A

Intron

14795

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15041	15041T>C	Intron
15343	15343G>A	Intron
15449	15449G>A	Intron
15502	15502G>A	Intron
15545	15545C>T	Intron
15589	15589A>G	Intron
15769	15769C>T	3'
15839	15839A>G	3'
16148	16148G>A	3'
16198	16198T>G	3'
16202	16202G>T	Intron
X59618 X59618		
reductase M2 polypept		no nibolideleotide
128	(-67) G>A	5'
189	(-6) T>G	5'
524	330C>G	Silent
1399	1205T>A	3'
1464	1270G>A	3'
1636	1442C>T	3'
1738	1544C>T	3'
2259	2065T>C	3 '
S72487 S72487		•
phosphorylase, partia		LD Thymidine
183	19G>A	D7N
483	319C>T	D7N
601		3'
1299	437G>C	3'
	1135G>A 131222 GEN-LUB	3'
		Thymidine
phosphorylase, promote 124		
439	124C>T	3'
1044	439G>A	3'
1331	1044^insCT	3'
1977	1331G>A	3'
2149	1977G>A	Intron
2467	2149G>A	Intron
2634	2467A>G	Intron
2975	2634C>G	Intron
	2975G>A	Intron
3116	3116G>T	Intron
3255	3255A>C	Intron
3344	3344T>C	Intron
4051	4051C>A	Intron
4782	4782G>A	Intron
5022	5022T>C	Intron
5266	5266G>A	Intron

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5285 5438 5482 5629 5648 5731 M98045 M98045 folylpolyglutamate synthe		Homo sapiens ete cds (SEQ ID NO:12)
1747	1677G>T	Silent 3'
1900	1830T>C	3 '
	510 GEN-LUE	Human
folylpolyglutamate synthe	tase (FPGS) gene	e. exons $5-11$. and
partial cds (SEQ ID NO:13)	, enems o 11, and
1424	1424C>A	Intron
1649	1649G>A	Intron
2554	2554A>G	Intron
U24252 U24252	136510 GEN-L	UF
Folylpolyglutamate synthe	tase, promoter a	nd exons 1-4 (SEQ ID
NO:14)		
263	263A>G	Intron
266	266G>T	Intron
527	527C>G	Intron
1037	1037A>G	5'
1139	1139G>A	Intron
1217 1647	1217C>T	Intron
1955	1647C>T	Intron
2017	1955G>A	Intron
2017	2017G>A	Intron
2189	2037G>A	Intron
2282	2189A>G	Intron
2309	2282C>T 2309A>G	Intron
U09806 U09806	236250 GEN-4:	Intron
methylenetetrahydrofolate		FZ Human
NO: 15)	reductase mixim,	parcial cds (SEQ ID
120	120T>C	Silent
464	464T>G	M155R
519	519C>T	Silent
668	668C>T	A223V
1059	1059T>C	Silent
1289	1289C>A	3'
1308	1308T>C	3'
1784	1784G>A	3'

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AF061655 deaminase,	AF061655 promoter	123920 (SEQ ID NO:	GEN-LUJ 16)	Cytidine
	575		575T>C	Intron
	648	•	648T>C	Intron
	771	•	771G>C	Intron
	883	{	383G>A	Intron
	941		l^insC	5'

1051A>C

K27Q

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